

Fig. 1A

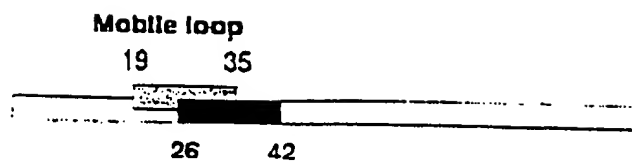


Fig. 1B



Fig. 2A

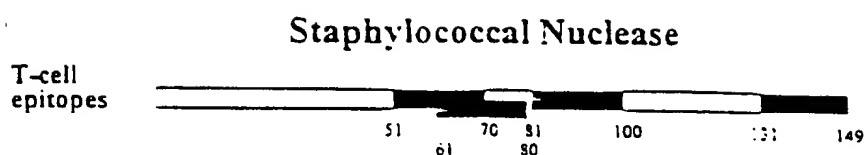
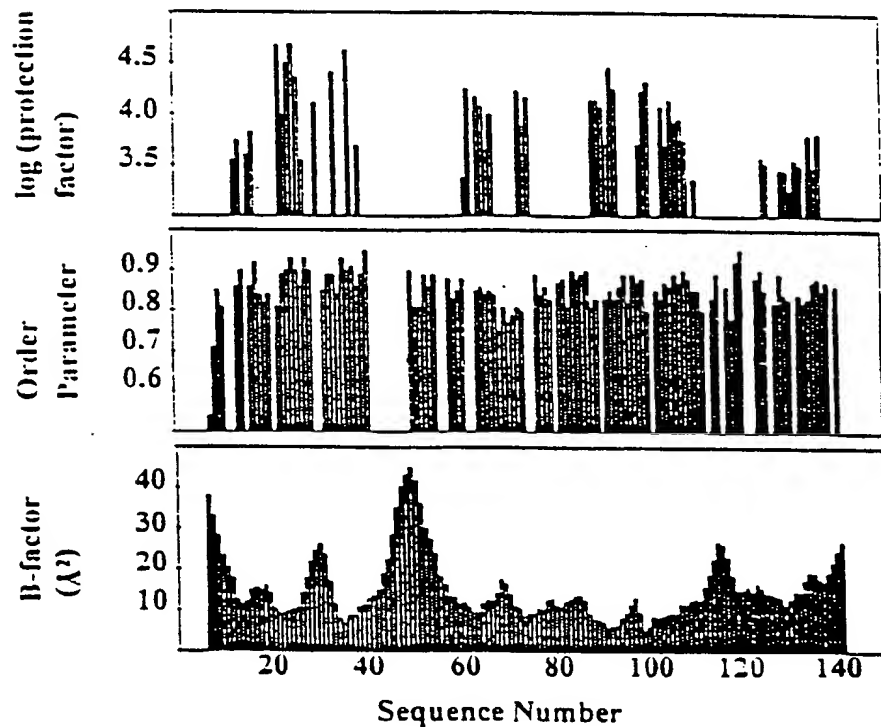


Fig. 2B



Lysozyme

Fig. 3A

T-cell
epitopes

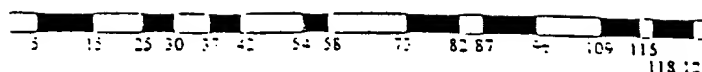
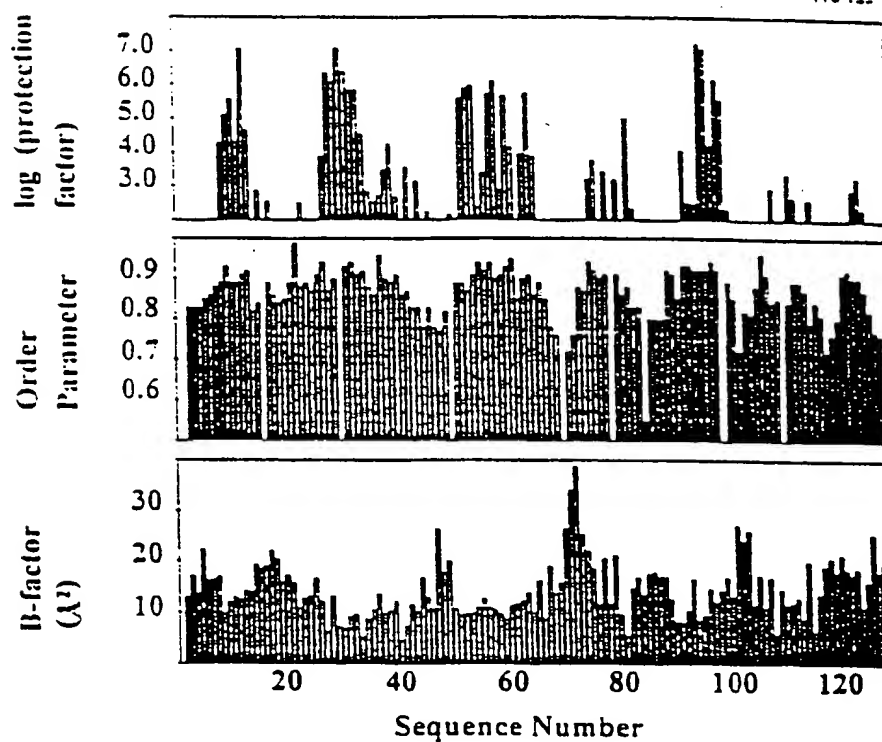
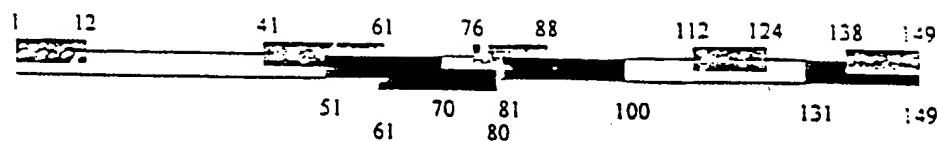


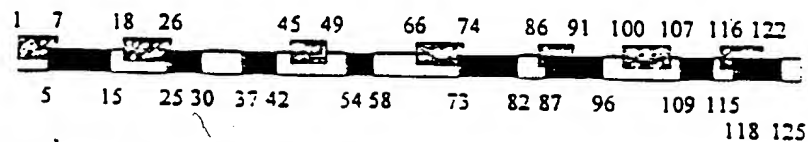
Fig. 3B



Staphylococcal Nuclease



Lysozyme



Cytochrome c

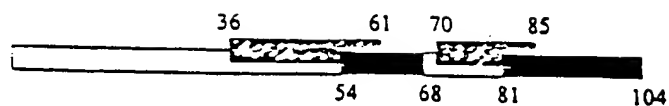


Figure 4

Fig. 5A

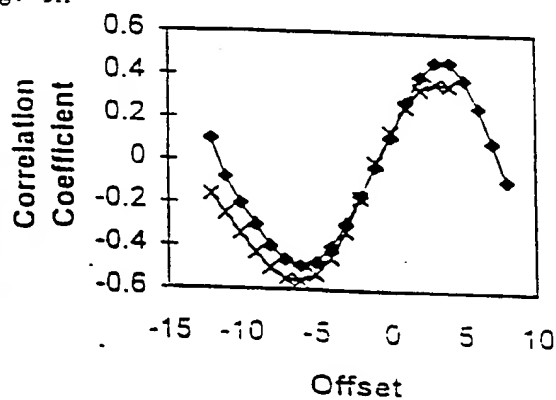
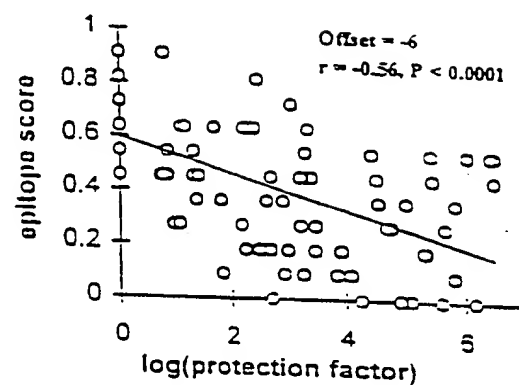


Fig. 5B



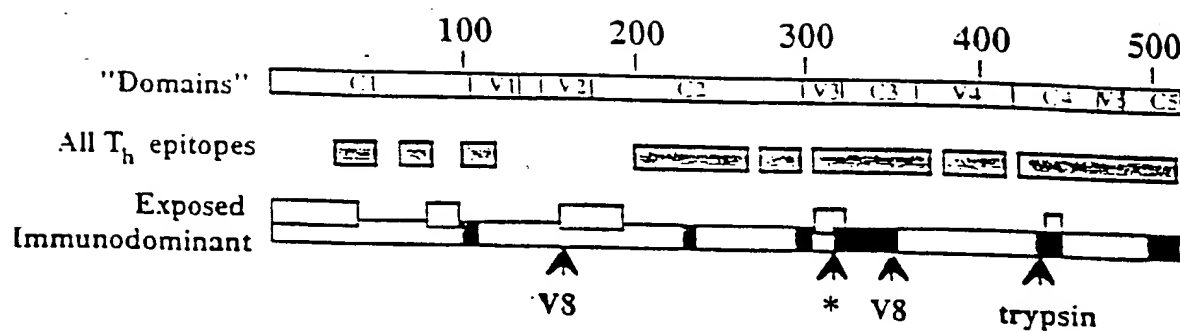
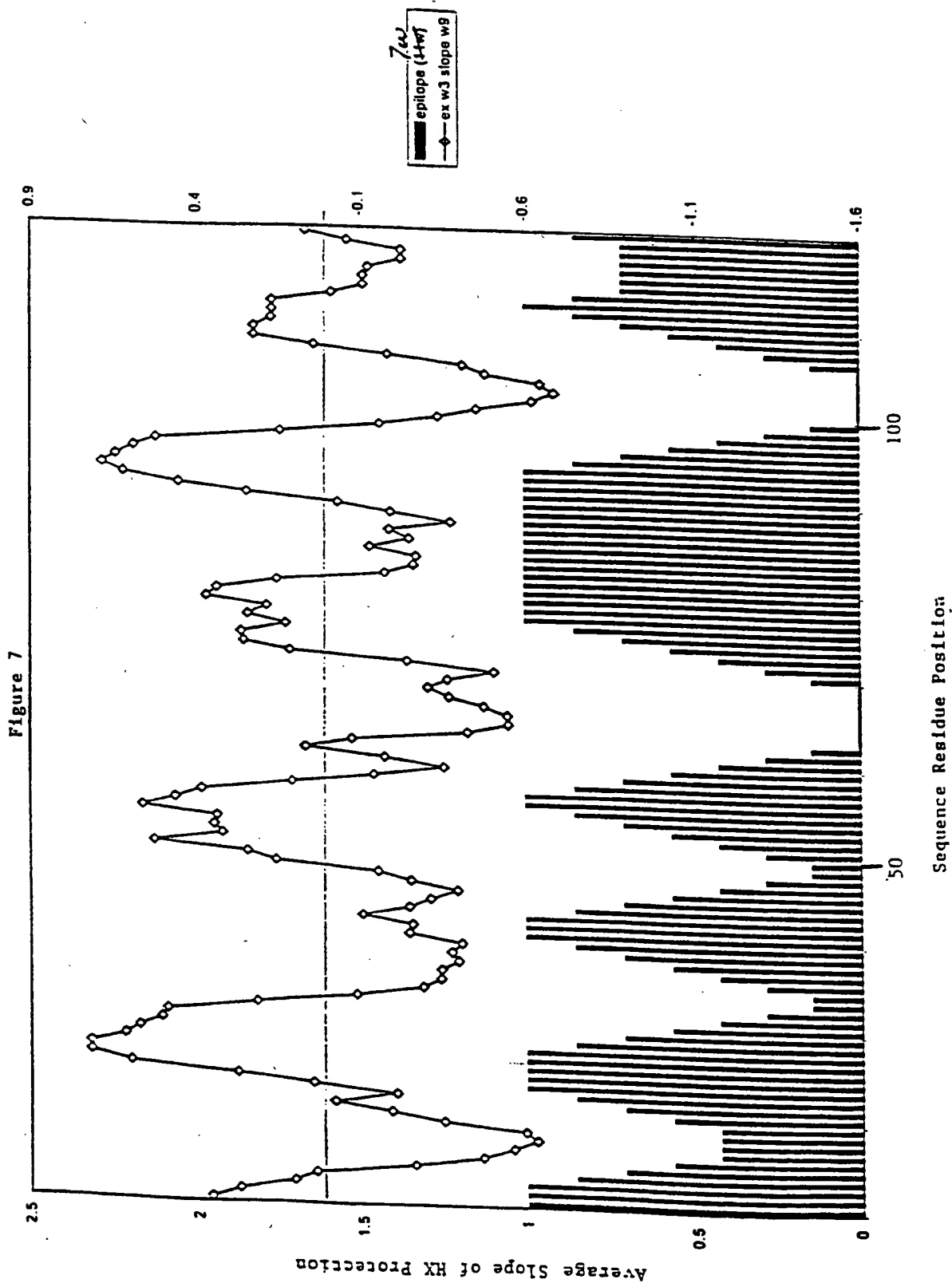


Figure 6



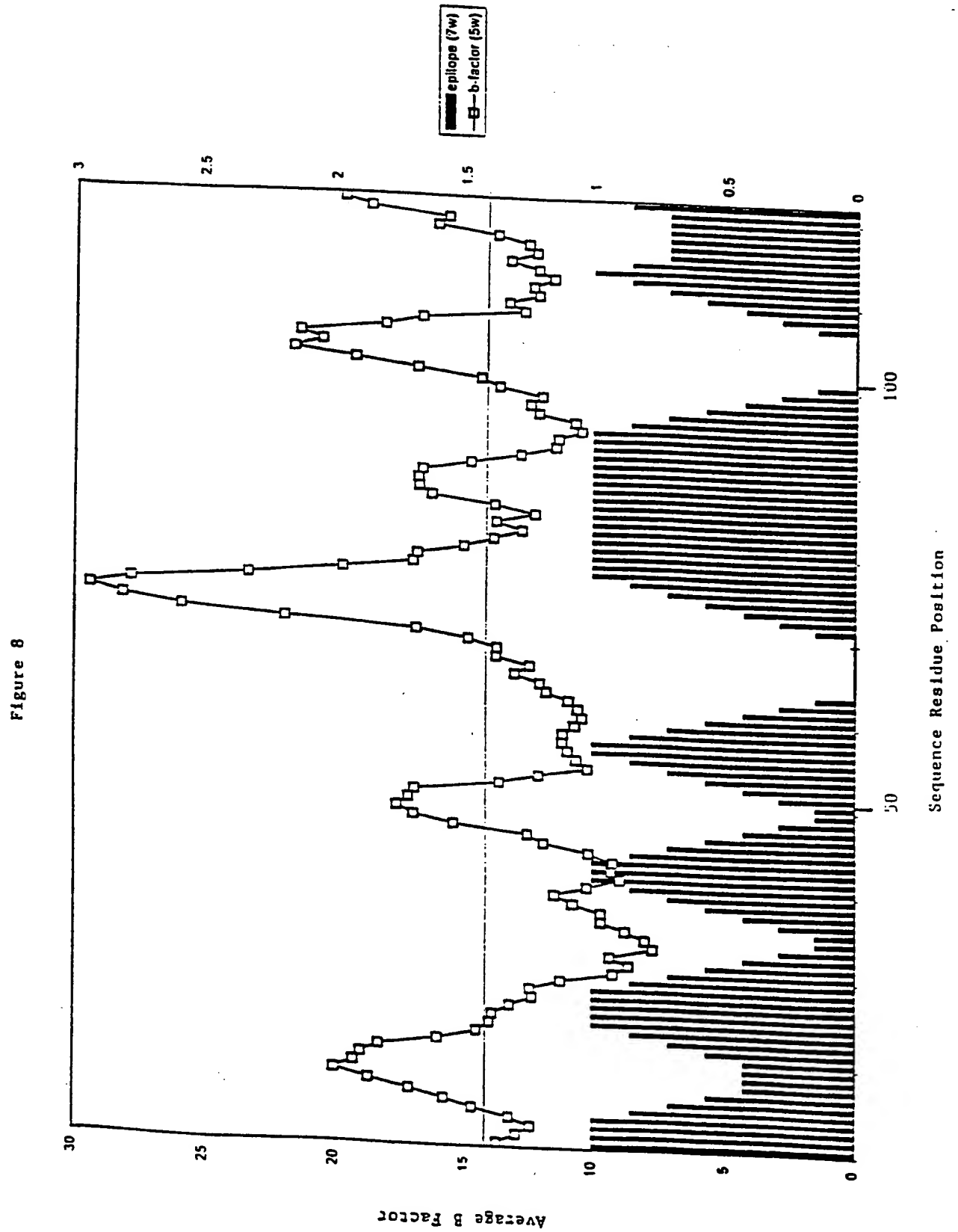


Figure 9

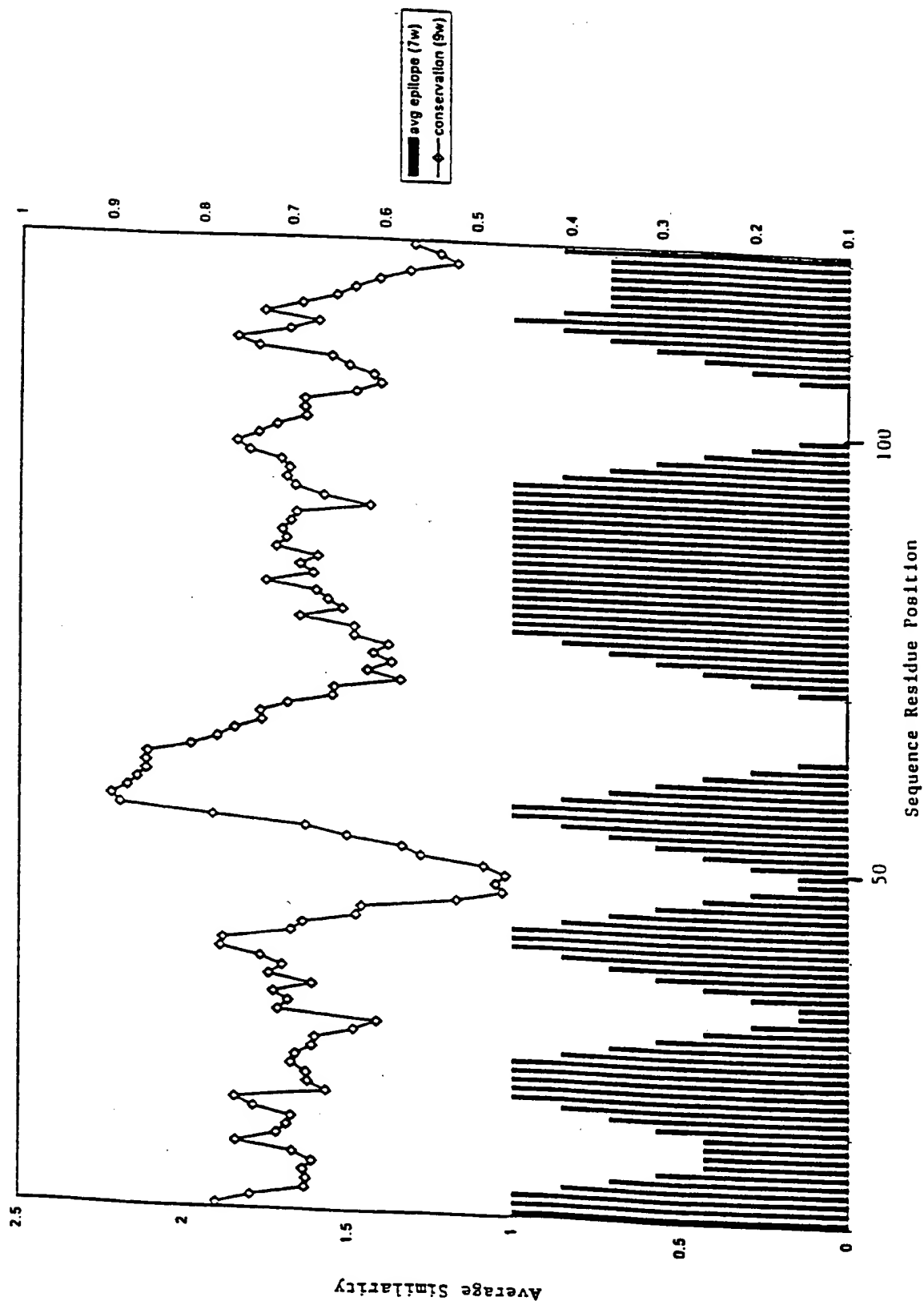
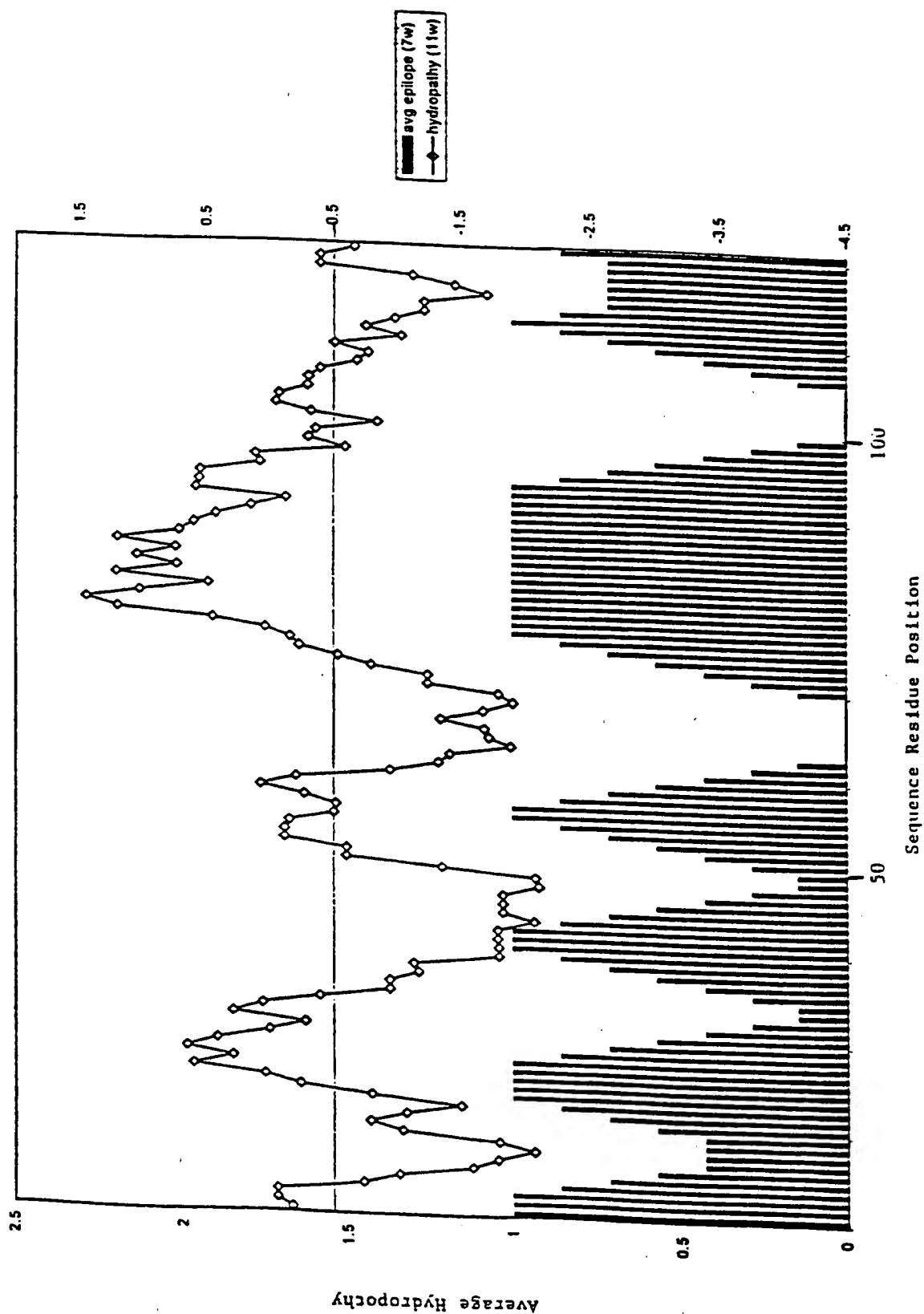


Figure 10



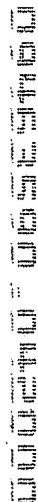
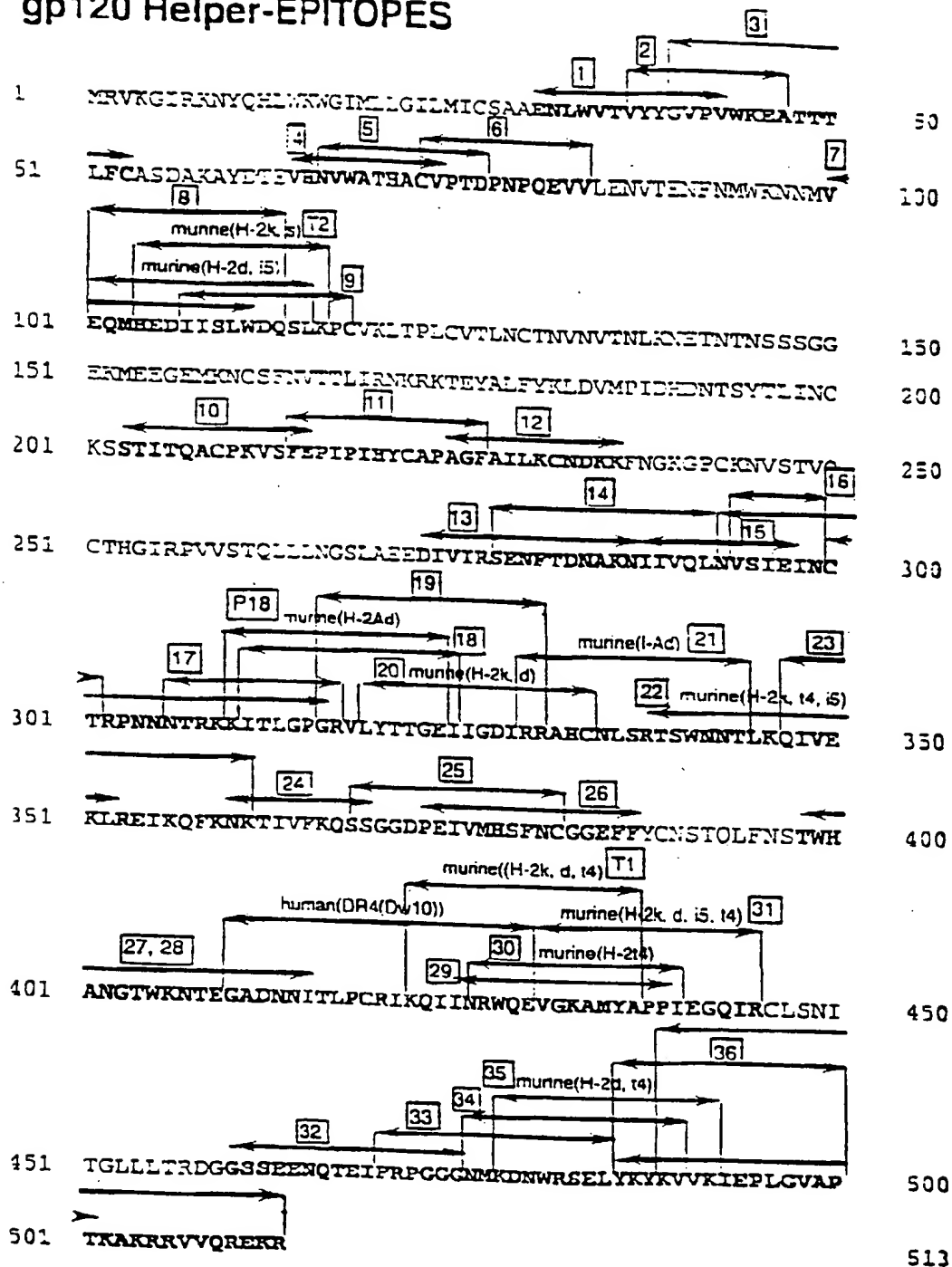


Figure 11A

Figure 11B

gp120 Helper-EPITOPES



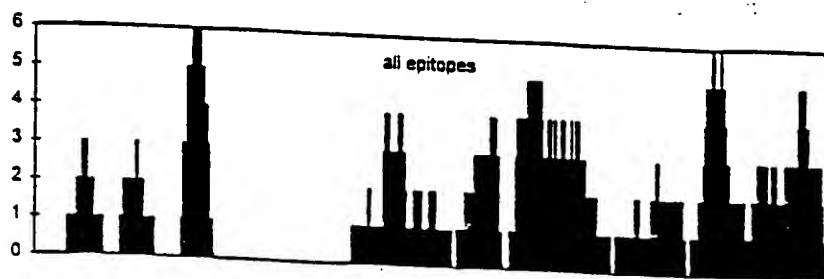


Figure 13A

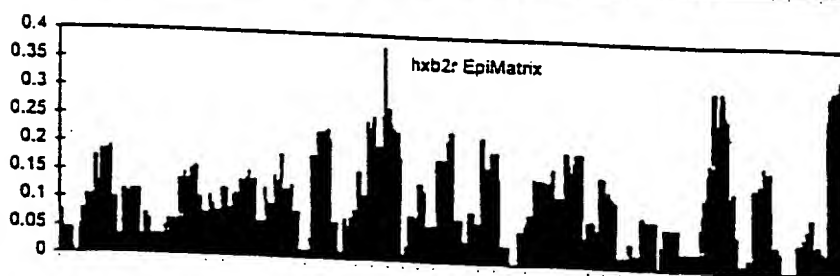


Figure 13B

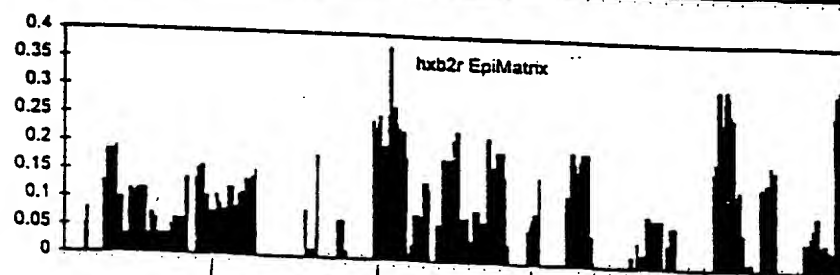


Figure 13C

Figure 1 consists of 12 histograms arranged in a 6x2 grid. The columns are labeled 'n = 10' and 'n = 20'. The rows are labeled 'm = 10', 'm = 20', 'm = 30', 'm = 40', 'm = 50', and 'm = 60'. The x-axis for all histograms is 'Number of non-zero elements' ranging from 0 to 100. The y-axis is 'Frequency' ranging from 0 to 10. The distributions are roughly bell-shaped and centered around 50 for n=10 and 100 for n=20.

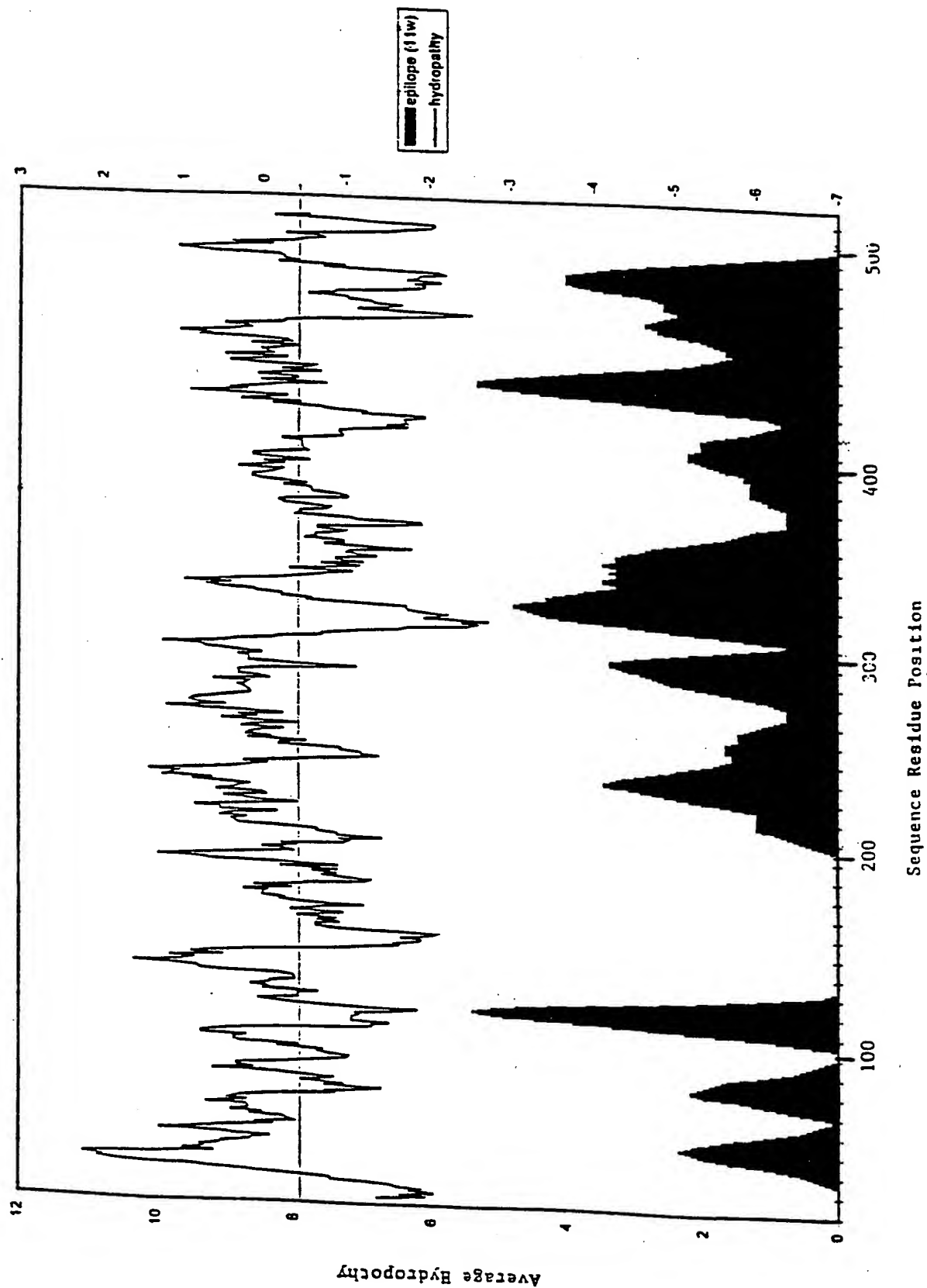
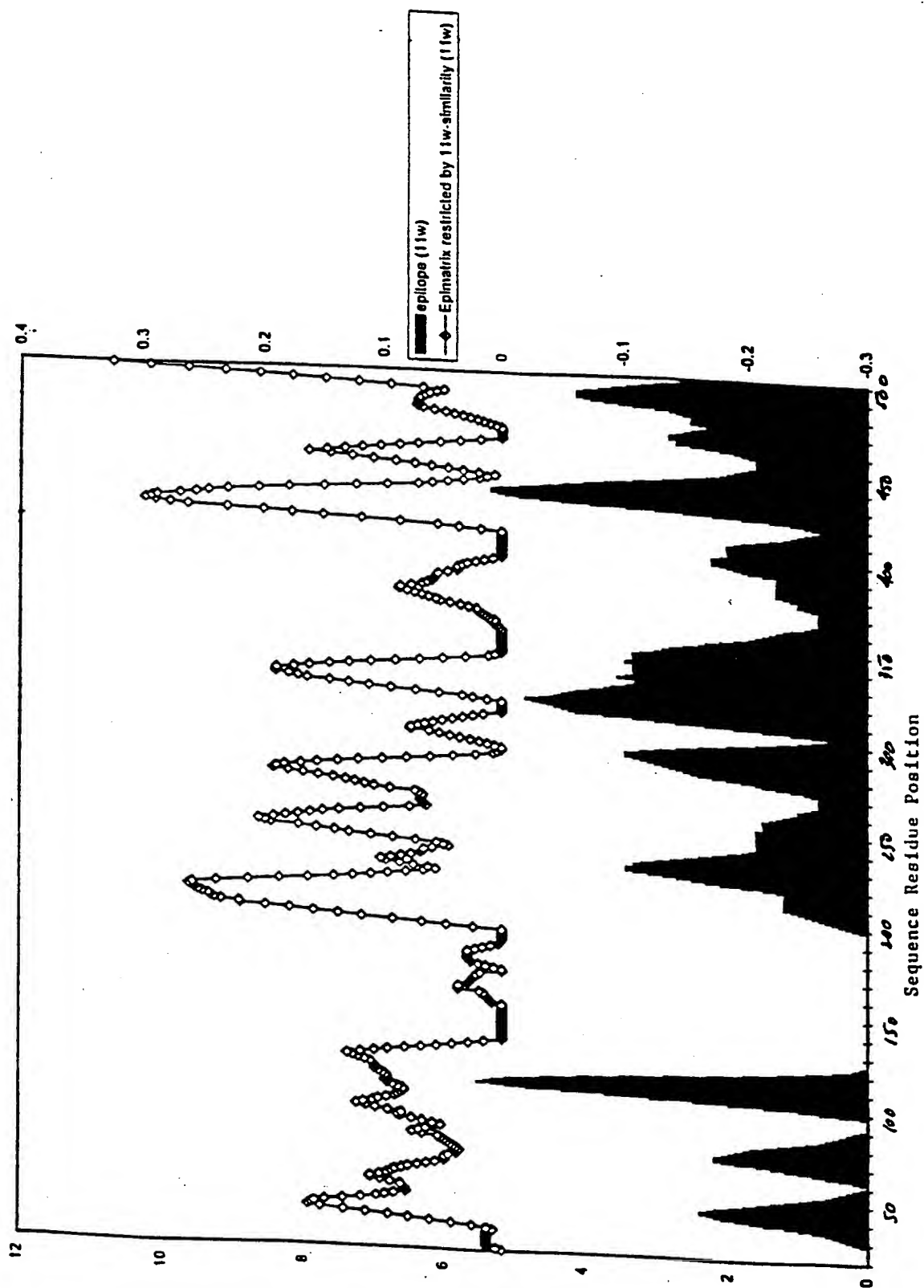


Figure 14



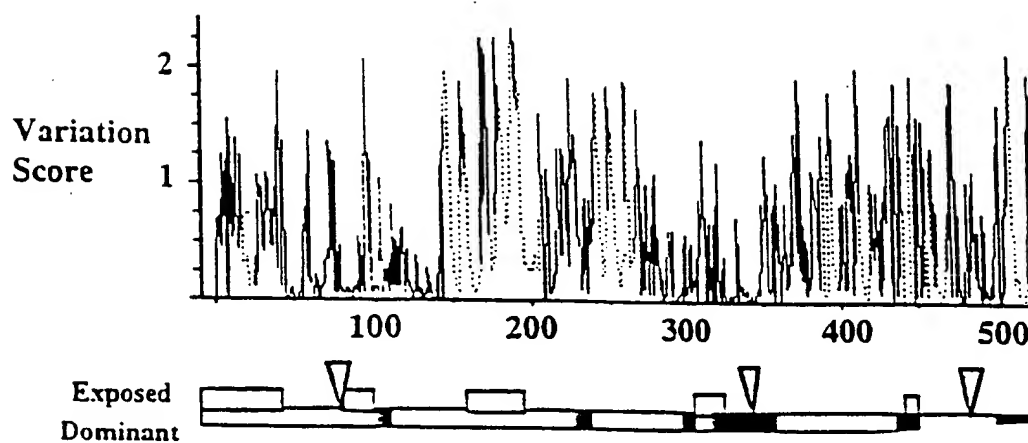


Figure 15

Fig. 16A

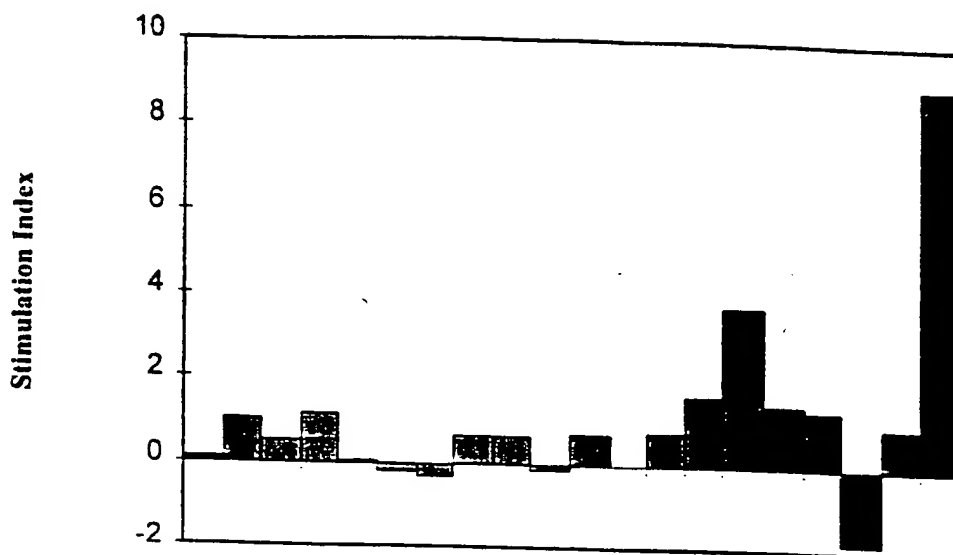


Fig. 16B

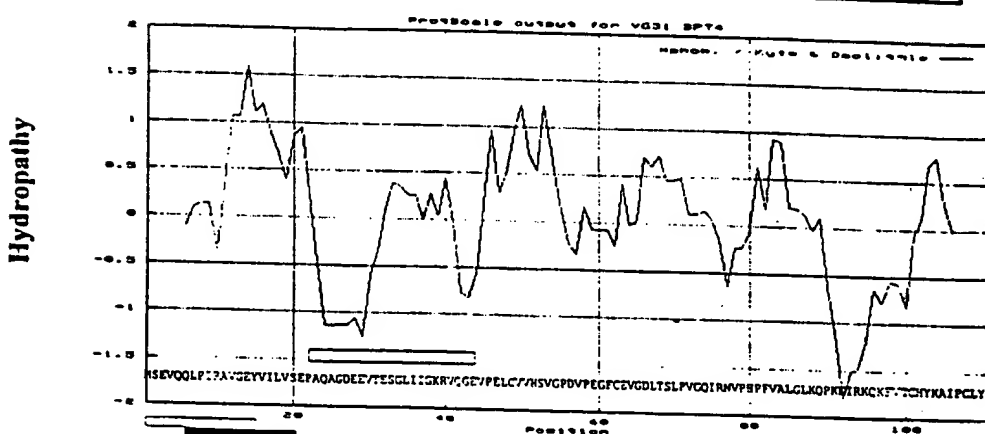


Fig. 16C

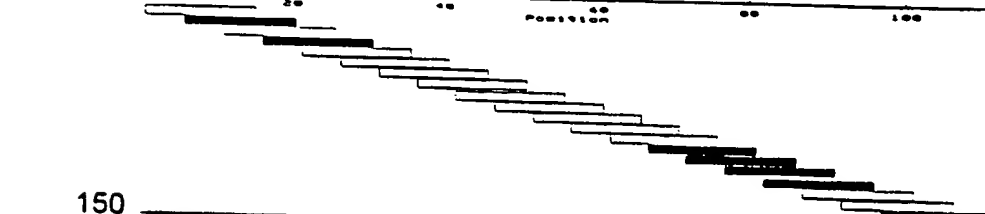


Fig. 16D

